>33358 cDNA ORF 75-1046

>33358 amino acids

MEDGPVFYGFKNIFITMFATFFFFKLLIKVFLALLTHFYIVKGNRKEAARIAEEIYGGISDCWADRSPLH EAAAQGRLLALKTLIAQGVNVNLVTINRVSSLHEACLGGHVACAKALLENGAHVNGVTVHGATPLFNACC SGSAACVNVLLEFGAKAQLEVHLASPIHEAVKRGHRECMEILLANNVNIDHEVPQLGTPLYVACTYQRVD CVKKLLELGASVDHGQWLDTPLHAAARQSNVEVIHLLTDYGANLKRRNAQGKSALDLAAPKSSVEQALLL REGPPALSQLCRLCVRKCLGRACHQAIHKLHLPEPLERFLLYQ.

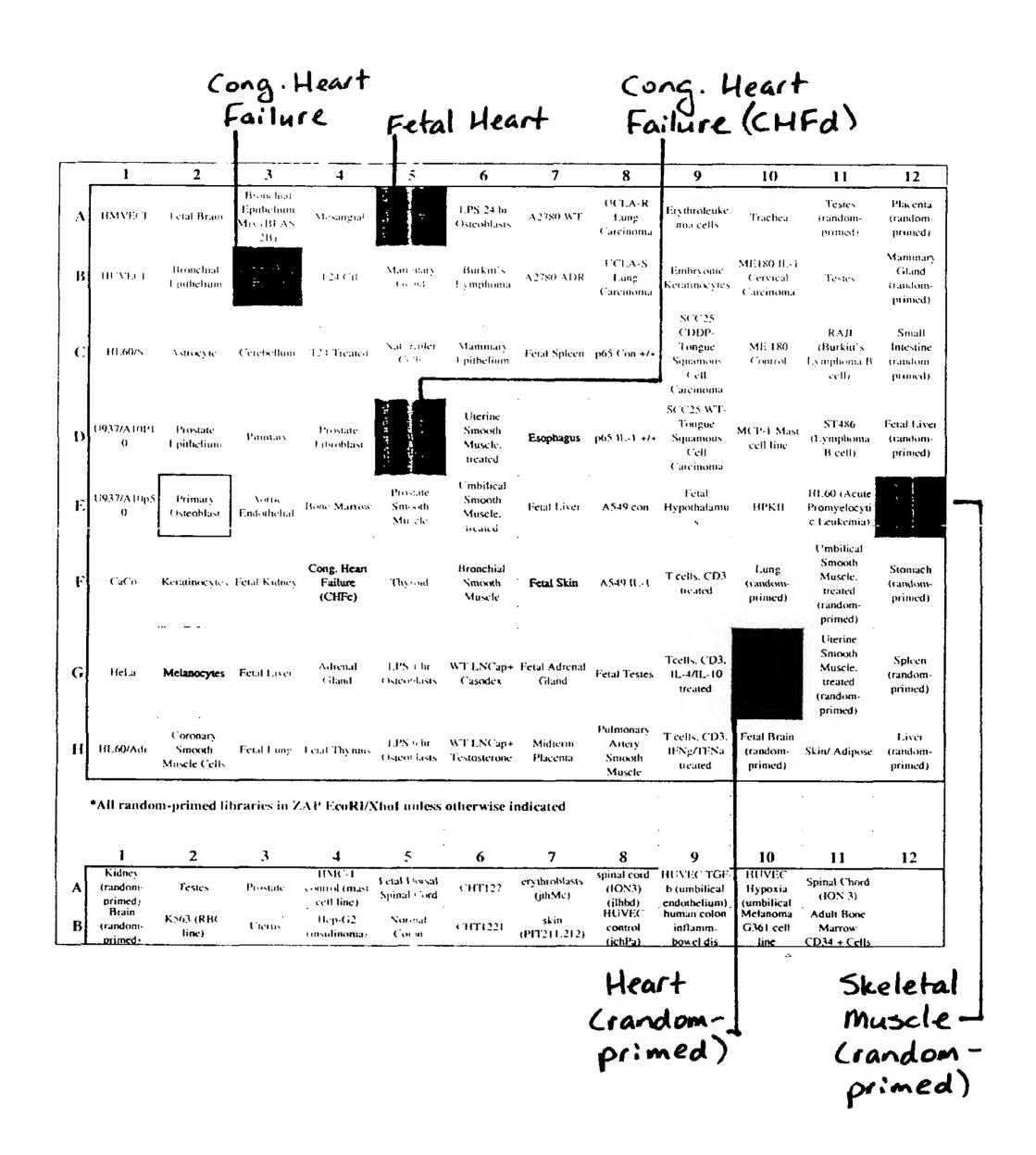
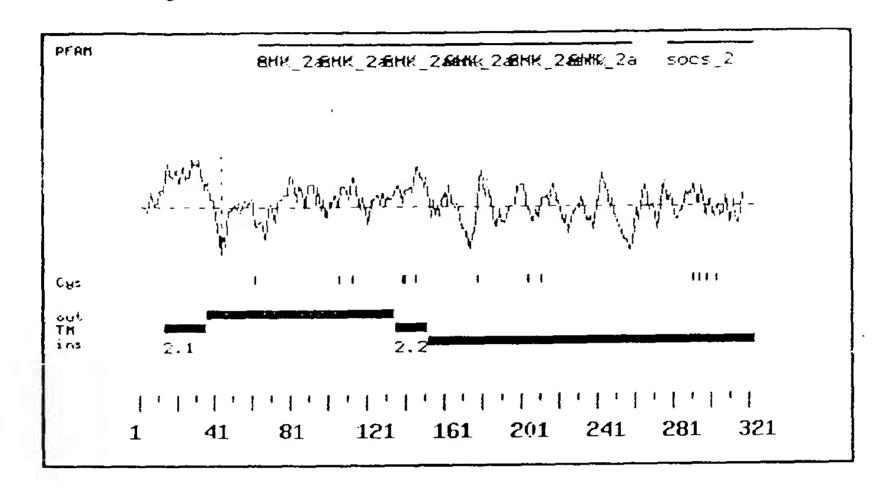


Figure 2A

		Hea	1+								
C	HPK (random- primed)	Lang	Mammary Cland	Normal Breasi Epithetia	Colon to Liver Metastases (CHT128)	normal breasi	BMCD34+ (jyhMc)	W138 20hi serum starve emb. Lung	Tht Cells	HUVEC untreated (umbilical endothelia)	
D	Cierus Gandom primed)	Liver	Spleen	Normal Ovarian Epithelia	Colon to Liver Metastasis (CHT133)	PTH Osteo	ovarian ascites(johOb)	hing squamous celf carcinoma	Th2 Cells	1BD Colon (WUM 23)	
E	Thymus trandom primed)	Be a	Small Intestruc	Normal Megakarocyt es	Colon Carcinoma (NDR109)	fung adenocarcino ma(PFF245)	(BI) colon (WUM6) (jbhna)	brain subcortical white matter	Prostate Tomot Xenografi A U	Trigeminal Ganglia	
F	9 week Fetus	Thymus	Retinal Pigmenosa Epithelia	Bone Marrow	Colon Carcinoma (NDR103)	lung squamous cell carcinoma PET299	Cervical Cancer (johCe)	normal prostate(ziplo x)	Prostate Tumor Xenograft K10	Lumbosacaral Spinal Chord	
G	A549 control (random- primed)	Stomach	Ketma	Th-Linduced Ticelt	Colon Carcinoma (NDR82)	d8 dendruic cells	Spinal cord (jlbbe) ~	ovarian epithelium tumor	Prostate Cancer to Liver Metastasis JHH3	Lumbosacaral Dorsal Root Ganglia	
н	Salivary Gland	Skeletał Musele	HMC(4 cmast cell line)	Th-2 induced Ticell	Colon Carcinoma (NDR097)	Megakaryocy tes(jtheb)	DRG (ION6.7.8) (jilixe)	nuvec Laname (umbilical end.)	Prostate Cancer to Liver Metastasis JHH4	Dorsal Root Ganglia (ION 6.7 &8)	

Analysis of 33358 (323 aa)



>33358
MEDGPVFYGFKNIFITMFATFFFKLLIKVFLALLTHFYIVKGNRKEAARIAEEIYGGIS
DCWADRSPLHEAAAQGRLLALKTLIAQGVNVNLVTINRVSSLHEACLGGHVACAKALLEN
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ILLANNVNIDHEVPQLGTPLYVACTYQRVDCVKKLLELGASVDHGQWLDTPLHAAARQSN
VEVIHLLTDYGANLKRRNAQGKSALDLAAPKSSVEQALLLREGPPALSQLCRLCVRKCLG
RACHQAIHKLHLPEPLERFLLYQ

FIGURE 3